

SEQUENCE LISTING

<110> MOECKEL, Bettina
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<120> NUCLEOTIDE SEQUENCES WHICH CODE FOR THE *rpoB* GENE

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<151> 2001-02-16

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<170> PatentIn version 3.0

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430

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 Asp Thr Ala
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Ser Trp Leu Ile Gly Thr Pro Glu Trp Arg Ala Arg Gln Lys Glu Glu
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 Phe Gly Phe Ser Glu Ile Met Met Ser Thr Leu Glu Ser Asp Gly Val
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 Gly Glu Gln Pro Thr Arg Asp Leu Ala Gln Ser Leu Leu Asp Asn Ser
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 Ile Asn Arg Lys Leu Gly Leu Gly Gly Asp His Asp Gly Leu Met Thr
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Leu Thr Glu Glu Asp Ile Ala Thr Thr Ile Glu Tyr Leu Val Arg Leu
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His Ala Gly Glu Arg Val Met Thr Ser Pro Asn Gly Glu Glu Ile Pro
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Val Glu Thr Asp Asp Ile Asp His Phe Gly Asn Arg Arg Leu Arg Thr
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Asn Ile Gly Leu Ile Gly Ser Leu Ala Ser Tyr Ala Arg Val Asn Pro
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570

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Val Ser Ala Asp Phe Ile Thr Ile Met Ala Asp Asp Gly Lys Arg Glu
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Thr Tyr Leu Leu Arg Lys Phe Gln Arg Thr Asn Gln Gly Thr Ser Tyr
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675 680 685

Val Ile Ala Asp Gly Pro Gly Thr Phe Asn Gly Glu Met Ser Leu Gly
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Arg Asn Leu Leu Val Ala Phe Met Pro Trp Glu Gly His Asn Tyr Glu
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<212> DNA

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Arg Val Thr Ser Gly Leu Glu Asn Ile Leu Glu Glu Leu Ser Pro Ile	
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Lys Ser Thr Glu Arg Pro Leu His Ala Val Lys Val Ile Pro Phe Arg	
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Gly Leu Gly Gly Asp His Asp Gly Leu Met Thr Leu Thr Glu Glu Asp	
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Ile Ala Thr Thr Ile Glu Tyr Leu Val Arg Leu His Ala Gly Glu Arg	
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211> 1165

212> PRT

213> Corynebacterium glutamicum

300> 4

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Ser Trp Leu Ile Gly Thr Pro Glu Trp Arg Ala Arg Gln Lys Glu Glu
 50 55 60

Phe Gly Glu Gly Ala Arg Val Thr Ser Gly Leu Glu Asn Ile Leu Glu
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Ser Glu Pro Arg Phe Glu Asp Val Lys Asn Thr Ile Asp Glu Ala Lys
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Glu Lys Asp Ile Asn Tyr Ala Ala Pro Leu Tyr Val Thr Ala Glu Phe
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Asp Phe Pro Met Met Thr Asp Lys Gly Thr Phe Ile Ile Asn Gly Thr
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Val Ile Pro Phe Arg Gly Ala Trp Leu Glu Phe Asp Val Asp Lys Arg
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Asp Ser Val Gly Val Arg Ile Asp Arg Lys Arg Arg Gln Pro Val Thr
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Thr Pro Thr Ser Leu Ile Asn Val Arg Pro Val Ser Ala Ala Ile Arg
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485 490 495

Phe Gly Phe Ile Glu Thr Pro Tyr Arg Arg Ile Ile Asp Gly Lys Leu
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Thr Asp Gln Ile Asp Tyr Leu Thr Ala Asp Glu Glu Asp Arg Phe Val
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565 570 575

Val Gly Thr Ala Met Ile Pro Phe Leu Glu His Asp Asp Ala Asn Arg
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Ala Leu Met Gly Ala Asn Met Gln Lys Gln Ala Val Pro Leu Ile Arg
595 600 605

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Val Ser Ala Asp Phe Ile Thr Ile Met Ala Asp Asp Gly Lys Arg Glu
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Thr Tyr Leu Leu Arg Lys Phe Gln Arg Thr Asn Gln Gly Thr Ser Tyr
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675 680 685

Val Ile Ala Asp Gly Pro Gly Thr Phe Asn Gly Glu Met Ser Leu Gly
690 695 700

Arg Asn Leu Leu Val Ala Phe Met Pro Trp Glu Gly His Asn Tyr Glu
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Asp Ala Ile Ile Leu Asn Gln Asn Ile Val Glu Gln Asp Ile Leu Thr
725 730 735

Ser Ile His Ile Glu Glu His Glu Ile Asp Ala Arg Asp Thr Lys Leu
740 745 750

Gly Ala Glu Glu Ile Thr Arg Asp Ile Pro Asn Val Ser Glu Glu Val
755 760 765

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770 775 780

Arg Asp Gly Asp Ile Leu Val Gly Lys Val Thr Pro Lys Gly Glu Thr
785 790 795 800

Glu Leu Thr Pro Glu Glu Arg Leu Leu Arg Ala Ile Phe Gly Glu Lys
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820 825 830

Gly Lys Val Ile Gly Val Arg His Phe Ser Arg Glu Asp Asp Asp Asp

835

840

845

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850 855 860

Arg Lys Ile Gln Asp Gly Asp Lys Leu Ala Gly Arg His Gly Asn Lys
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885 890 895

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900 905 910

Arg Met Asn Ile Gly Gln Val Leu Glu Thr His Leu Gly Trp Leu Ala
915 920 925

Ser Ala Gly Trp Ser Val Asp Pro Glu Asp Pro Glu Asn Ala Glu Leu
930 935 940

Val Lys Thr Leu Pro Ala Asp Leu Leu Glu Val Pro Ala Gly Ser Leu
945 950 955 960

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965 970 975

Leu Leu Ala Asn Ser Arg Pro Asn Arg Asp Gly Asp Val Met Val Asn
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Ala Asp Gly Lys Ala Thr Leu Ile Asp Gly Arg Ser Gly Glu Pro Tyr
995 1000 1005

Pro Tyr Pro Val Ser Ile Gly Tyr Met Tyr Met Leu Lys Leu His
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His Leu Val Asp Glu Lys Ile His Ala Arg Ser Thr Gly Pro Tyr
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211> 5099
212> DNA
213> Corynebacterium glutamicum

220>
221> CDS
222> (702)..(4196)

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Val Leu Glu Gly Pro	5
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Ile Leu Ala Val Ser Arg Gln Thr Lys Ser Val Val Asp Ile Pro Gly	20
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Ala Pro Gln Arg Tyr Ser Phe Ala Lys Val Ser Ala Pro Ile Glu Val	35
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Pro Gly Leu Leu Asp Leu Gln Leu Asp Ser Tyr Ser Trp Leu Ile Gly	50
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Thr Pro Glu Trp Arg Ala Arg Gln Lys Glu Glu Phe Gly Glu Gly Ala	65
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Arg Val Thr Ser Gly Leu Glu Asn Ile Leu Glu Leu Ser Pro Ile	85
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Gln Asp Tyr Ser Gly Asn Met Ser Leu Ser Leu Ser Glu Pro Arg Phe	100
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Glu Asp Val Lys Asn Thr Ile Asp Glu Ala Lys Glu Lys Asp Ile Asn	115
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Gly Ala Trp Leu Glu Phe Asp Val Asp Lys Arg Asp Ser Val Gly Val	210
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Arg Ile Asp Arg Lys Arg Arg Gln Pro Val Thr Val Leu Leu Lys Ala	225
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gca ttg ctg gag atc tac cgc aag cag cgt cca ggc gag cag cct acc Ala Leu Leu Glu Ile Tyr Arg Lys Gln Arg Pro Gly Glu Gln Pro Thr 265 270 275	1532
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acc cca tac cgt cgc atc atc gac ggc aag ctg acc gac cag att gac				2252
Thr Pro Tyr Arg Arg Ile Ile Asp Gly Lys Leu Thr Asp Gln Ile Asp	505	510	515	
tac ctt acc gct gat gag gaa gac cgc ttc gtt gtt gcg cag gca aac				2300
Tyr Leu Thr Ala Asp Glu Glu Asp Arg Phe Val Val Ala Gln Ala Asn	520	525	530	
acg cac tac gac gaa gag ggc aac atc acc gat gag acc gtc act gtt				2348
Thr His Tyr Asp Glu Gly Asn Ile Thr Asp Glu Thr Val Thr Val	535	540	545	
cgt ctg aag gac ggc gac atc gcc atg gtt ggc cgc aac gcg gtt gat				2396
Arg Leu Lys Asp Gly Asp Ile Ala Met Val Gly Arg Asn Ala Val Asp	550	555	560	565
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Tyr Met Asp Val Ser Pro Arg Gln Met Val Ser Val Gly Thr Ala Met	570	575	580	
att cca ttc ctg gag cac gac gat gct aac cgt gca ctg atg ggc gcg				2492
Ile Pro Phe Leu Glu His Asp Asp Ala Asn Arg Ala Leu Met Gly Ala	585	590	595	
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Asn Met Gln Lys Gln Ala Val Pro Leu Ile Arg Ala Glu Ala Pro Phe	600	605	610	
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Val Gly Thr Gly Met Glu Gln Arg Ala Ala Tyr Asp Ala Gly Asp Leu	615	620	625	
gtt att acc cca gtc gca ggt gtg gtg gaa aac gtt tca gct gac ttc				2636
Val Ile Thr Pro Val Ala Gly Val Val Glu Asn Val Ser Ala Asp Phe	630	635	640	645
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Ile Thr Ile Met Ala Asp Asp Gly Lys Arg Glu Thr Tyr Leu Leu Arg	650	655	660	
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Lys Phe Gln Arg Thr Asn Gln Gly Thr Ser Tyr Asn Gln Lys Pro Leu	665	670	675	
gtt aac ttg ggc gag cgc gtt gaa gct ggc cag gtt att gct gat ggt				2780
Val Asn Leu Gly Glu Arg Val Glu Ala Gly Gln Val Ile Ala Asp Gly	680	685	690	
cca ggt acc ttc aat ggt gaa atg tcc ctt ggc cgt aac ctt ctg gtt				2828
Pro Gly Thr Phe Asn Gly Glu Met Ser Leu Gly Arg Asn Leu Leu Val	695	700	705	
gcg ttc atg cct tgg gaa ggc cac aac tac gag gat gcg atc atc ctc				2876
Ala Phe Met Pro Trp Glu Gly His Asn Tyr Glu Asp Ala Ile Ile Leu	710	715	720	725

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212> PRT

213> Corynebacterium glutamicum

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610

615

620

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850 855 860

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Asp Gly Thr Pro Val Asp Ile Ile Leu Asn Thr His Gly Val Pro Arg
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Pro Gln Arg Arg Gly Val Cys Thr Arg Val Tyr Thr Thr Thr Pro Lys

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Lys Pro Asn Ser Ala Leu	Arg Lys Val Ala Arg	Val Arg Leu Thr Ser	
45	50	55	
ggc atc gag gtt tcc gct	tac atc cct ggt gag	ggc cac aac ctg cag	724
Gly Ile Glu Val Ser Ala	Tyr Ile Pro Gly Glu	Gly His Asn Leu Gln	
60	65	70	75
gag cac tcc atg gtg ctc	gtt cgc ggt ggt cgt	gtt aag gac ctc cca	772
Glu His Ser Met Val Leu	Val Arg Gly Gly Arg	Val Lys Asp Leu Pro	
80	85	90	
ggt gtc cgt tac aag atc	gtc cgt ggc gca ctg	gat acc cag ggt gtt	820
Gly Val Arg Tyr Lys Ile	Val Arg Gly Ala Leu	Asp Thr Gln Gly Val	
95	100	105	
aag gac cgc aag cag gct	cgt tcc ccg cta cgg	cgc gaa gag ggg ata	868
Lys Asp Arg Lys Gln Ala	Arg Ser Pro Leu Arg	Arg Glu Gly Ile	
110	115	120	
att aaa aat gcg taaatcagca	gctcctaagc gtccagtagt	tcaggaccc	920
File Lys Asn Ala			
125			
gtatacaagt ccgagctcgt	taccagctc gtaacaaga	tcctcatcgg tggcaagaag	980
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<210> 8

<211> 127

<212> PRT

<213> Corynebacterium glutamicum

<400> 8

Met Pro Thr Ile Gln Gln Leu Val Arg Lys Gly Arg His Asp Lys Ser
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Ala Lys Val Ala Thr Ala Ala Leu Lys Gly Ser Pro Gln Arg Arg Gly
20 25 30

Val Cys Thr Arg Val Tyr Thr Thr Thr Pro Lys Lys Pro Asn Ser Ala
35 40 45

Leu Arg Lys Val Ala Arg Val Arg Leu Thr Ser Gly Ile Glu Val Ser
50 55 60

Ala Tyr Ile Pro Gly Glu Gly His Asn Leu Gln Glu His Ser Met Val
65 70 75 80

Leu Val Arg Gly Gly Arg Val Lys Asp Leu Pro Gly Val Arg Tyr Lys
85 90 95

Ile Val Arg Gly Ala Leu Asp Thr Gln Gly Val Lys Asp Arg Lys Gln
100 105 110

Ala Arg Ser Pro Leu Arg Arg Glu Glu Gly Ile Ile Lys Asn Ala
115 120 125